



## SEQUENCE LISTING

<110> University of Wales, Bangor

## <120> Schistosomiasis vaccine

<130> L213

<140>10/020,441

<141>2001-12-18

<140>09/413,810

<141>1999-10/07

<150> GB 9821821.7

<151> 1998-10-07

<160> 3

<170> PatentIn Ver. 2.1

<210> 1

<211> 1489

<212> DNA

<212> BRI

<220>

<223> Description of Artificial Sequence: dna

<400> 1

gtgtcaacct ggttgatacg tagtggtaaa cctgtgcac accgcactgaa attcccgttc 60  
atcgcatctt taacgacaga gagaacaatg tttacagstt cactagtctc aacgagagca 120  
gtactcacag ctggtcattt tggttgcctt ccattgccag tgattcgggt aagagatcga 180  
ctg0aaacac attgttgcctt aatgtaatcg attgatttca cagacagtgt ttgcgtgtc 240  
gtgcgttgtt gtttgttat gcagtgggtt tgcatacgatt ttaatgtcta tataacttgtt 300  
cttatttcag gtttcatttc tcacacttag gaat00ggcg accaacaagg catccatcac 360  
caaccgtctg gagttaaagg ggcaccagga tacatg0ccc tcttgttatgt cggcacgaca 420  
gaggagacca atcgacacaa cactcagtgg attcgat0at tgcaattgtt atgctggctc 480  
aaatggtcaa cttacagagt ggaatcagag tgcatacgat0c tgccacagcc atcgatatac 540  
ccgccccctg gaactgggtt ttccattgtt ggttatggaa0 agggatgata acgaccgtga 600  
tccgtcacgt aagaatgggtt gaatattgaa gaaaagttagt t0gttggta ataaacgaca 660  
tgactcagtc gtcaagtccaa tgcaggatcg ttattctgtt tgcaggatgt tctgtttgtt 720  
t0gtctgtct gtctacacttga tccgggtt gttatggta gggccctgtat aataacaact 780  
gtgttggat gactttgttga cagttcagta gcagagtgtt ttccatctcg gtcattgtt 840  
tggtgagggtt aggtgacgtt atgtgagggtt agttgaggta gattggatgg gatggatgt 900  
gatgtgatgg gatgatttggt accacttggaa ggagagaaga ctcatgaaat atctatgca0 960  
aacgatggaa gtgtgttggat tacatgaaatg aggggggtcaa tgcaggatgtt gatgtgttt 1020  
gagagtgggtt agatggaga0 gtgacttggat cgtcgaataat agtgcacatg tgattgtatg 1080  
tggactattt ttgtgtgggtt agtgtaaagg gtggatatttgc tgcaggatgtt tattttcgaa 1140  
attcacttgtt gtgtttgtt ttgttccgtt gtgtttgtt ctttctcattc tgcaggatgtt 1200  
gtgttgttac tgcatacgatgg tttgtgttgc tttgttgcac cacagggtcg agcgactata 1260  
atggaatgcc gacatgcgac caatggcaat cctatatgtt tgaaaggcagg tcagaatttc 1320  
ggacagttac cccgtccagg tgcaggatgtt ggacacttcc tcccatccct tcaagggtcca 1380  
gtactcggtt tgcgtatcaca tgggtgtcaca ctgcctaaacc ttcccgatat cattgtcgag 1440  
tatgccat0 gtggcttagaa tgggttggat tgcaggatgtt aatatttgc 1489

<210> 2

<211> 106

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:protein

<400> 2

Val Ser Phe Leu Thr Leu Arg Asn Gly Asp Gln Gln Gly Ile His His  
1 5 10 15

Gln Pro Ser Gly Val Lys Val Ala Pro Gly Tyr Met Pro Ser Cys Met  
20 25 30

Ser Ala Arg Gln Arg Arg Pro Ile Ala Gln Thr Leu Ser Gly Phe Asp  
35 40 45

Ile Ala Ile Val Met Leu Ala Gln Met Val Asn Leu Gln Ser Gly Ile  
50 55 60

Arg Val Ile Ser Leu Pro Gln Pro Ser Asp Ile Pro Pro Pro Gly Thr  
65 70 75 80

Gly Val Phe Ile Val Gly Tyr Gly Arg Asp Asp Asn Asp Arg Asp Pro  
85 90 95

Ser Arg Lys Asn Gly Gly Ile Leu Lys Lys  
100 105

<210> 3

<211> 16

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:protein

<400> 3

Val Gly Tyr Gly Arg Asp Asp Asn Asp Arg Asp Pro Ser Arg Lys Asn  
1 5 10 15

---